

ANNOTATED SHEET

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+1	TCCTAGTACCATAATATTACAAAACACACACCTCAGAAACGAAGCTTGC
+53	ACATAATGTCTCTTCAACAAGTACCCACAACAGAAACAACCACCAGGAAGAA
1	M S L Q Q V P T T E T T T R K N
+105	CCGAAACGAGGGGCGGAAACGATTTACCGACAAACAAATAAGTTTCCTAGAG
17	R N E G R K R F T D K Q I S F L E
+157	TACATGTTTGTAGACACAGTCGAGACCCGAGTTAAGGATGAAACACCAGTTGG
34	Y M F E T Q S R P E L R M K H Q L
+209	CACATAAACTCGGGCTTCATCCTCGTCAAGTGGCGATATGGTTCCAGAACAA
51	A H K L G L H P R Q V A I W F Q N K
+261	ACGCGCGCGATCAAAGTCGAGGCAGATTGAGCAAGAGTATAACGCGCTAAAG
69	R A R S K S R Q I E Q E Y N A L K
+313	CATAACTACGAGACGCTTGCGTCTAAATCCGAGTCTCTAAAGAAAGAGAATC
86	H N Y E T L A S K S E S L K K E N
+365	AGGCCCTACTCAATCAGgtatggttgcaaacttacaatgttgcatccaacta
103	Q A L L N Q
+417	tttaagtagttttgaattttgtgacaataaagattgacaaatgttggtttga
+469	taattgattaacagTTGGAGGTGCTGAGAAATGTAGCAGAAAAGCATCAAGA
109	L E V L R N V A E K H Q E
+521	GAAAACTAGTAGTAGTGGCAGCGGTGAAGAATCGGATGATCGGTTTACGAAC
122	K T S S S G S G E E S D D R F T N
+573	TCTCCGGACGTTATGTTTGGTCAAGAAATGAATGTTCCGTTTTCGCGACGGTT
139	S P D V M F G Q E M N V P F C D G
+625	TTGCGTACTTTGAAGAAGGAAACAGTTTGTGAGATTGAAGAACAACCTGCC
156	F A Y F E E G N S L L E I E E Q L P
+677	AGACCCTCAAAAGTGGTGGGAGTTCTAAAGAGTAAAGAAGGATGTAGAAGTA
174	D P Q K W W E F *
+729	GTAGAGTAAAACTAAACATACCAGATAGTTGGTTTACACTTTGT
Athb-1	LPEKRRRLTTEQVHLLKSFETENKLEPERKTQLAKKLGLPQVAVWFQNRARWTKQLERDYDLKSTYDQLLSNYDSIVMDNDKLRSEVTSLTEK
Athb-6	LSEKRRRLSINQVKALEKNFELNKLPERKVKLAQELGLQPROVAVWFQNRARWTKQLEKDYGVLTQYDSLRRHFDLSRRDNESLLQEISKLTKE
Athb-7	NKNNQRRFSDQIKSLEMMFSETRLEPRKKVQLARELGLQPROVAIWQNKRRWKSQLETEYNILRQNYDNLASQFESLKKEKQALVSELQRLKEA
Athb-12	KSNNQKRFNEEQIKSLELIFESETRLEPRKKVQVARELGLQPMQTIWFQNKRRWTKQLEKEYNTLRANYNNLASQFEIMKKEKQSLVSELQRLNEE
Hahb-4	RNEGRKRFTDKQISFLEYMFETQSRPELRMKHQLAHKLGHPQVAIWQNKRRSKSRQIEQYNAKHNKETLASKSESLLKKNQALLNQLVLRNV

FIG. 1

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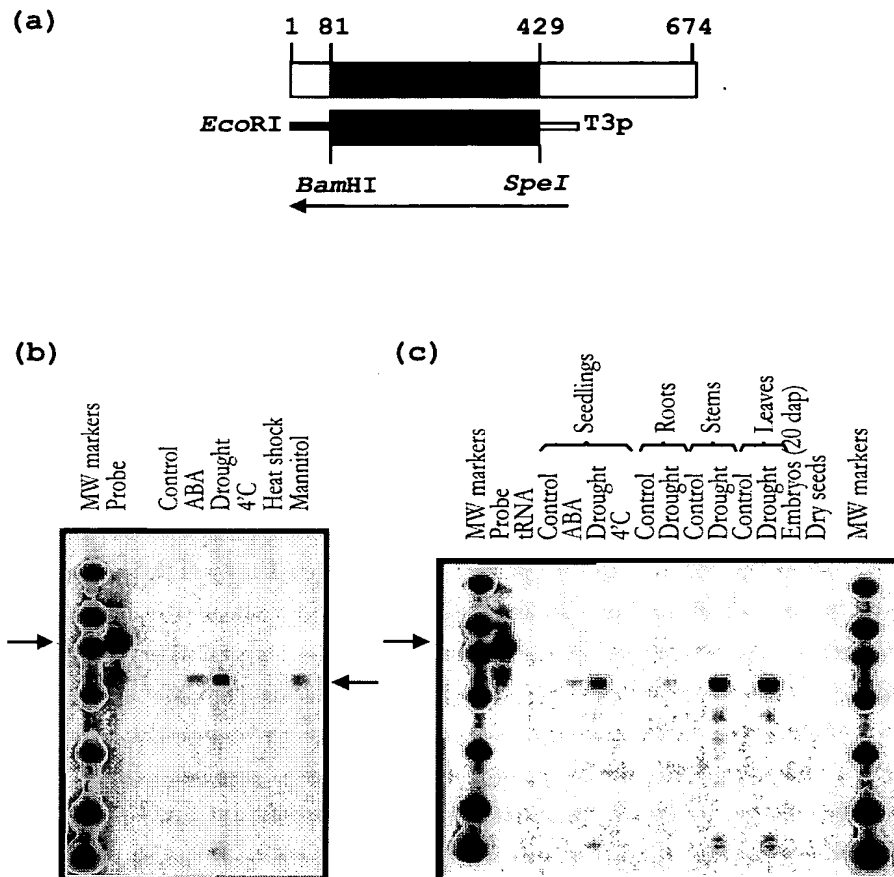


FIG. 2

ANNOTATED SHEET



FIG. 22

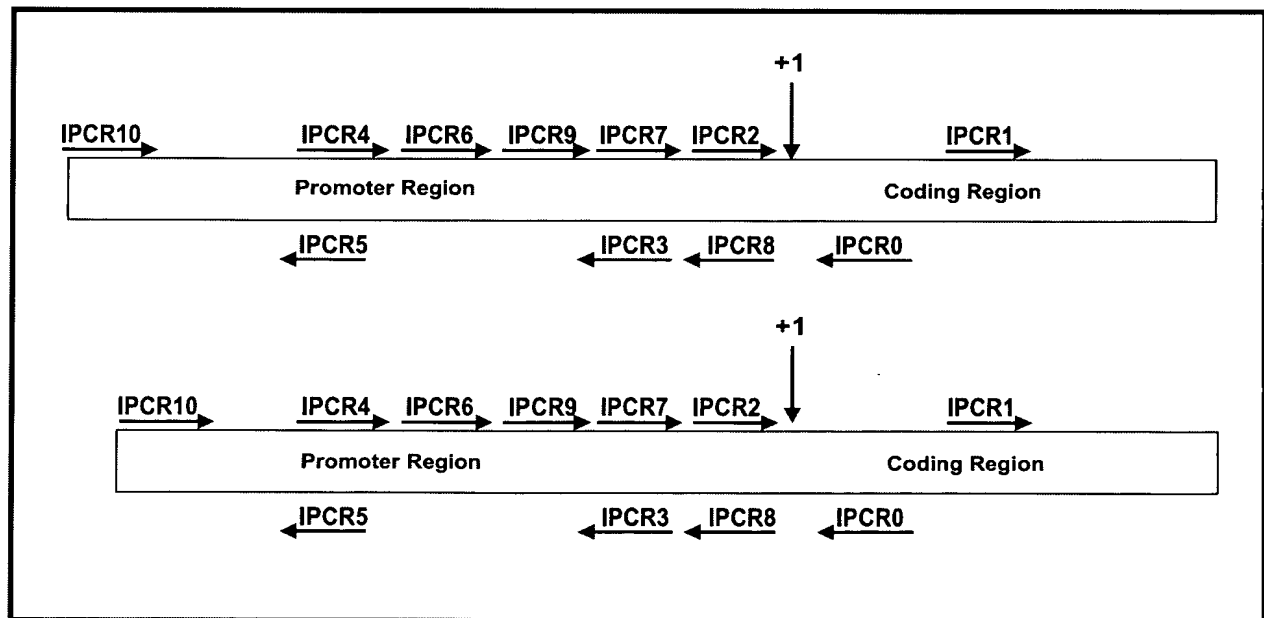


FIG. 23

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